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61 VYPGITEDQLWRAKYYDSAFHPDTGEKVVLLGRASAQVPMMTITGCALTFYRKTPTV 120
                                                                                                                                                                                                                                The present invention provides the protein and coding sequences for 43 novel human transport proteins (designated TPFTs). These can be used in the diagnosis and treament of transport, metabolic, neurological, reproductive, cardiovascular and immune disorders, and cell proliferative disorders such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 VFWQWVNQSFNAIVNYSNRSGDTPIIVRQLGTAYVSATTGAVATALGLKSLTKRLPPLVG 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         181 RPVPPAAVAANCINIPLARQRELQVGIPVADEAGQRLGYSVTAAKQGIPQVVISRICMA 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          191 RFVPFAANAAANCINIPLARQRELQVGIPVADEAGQRLGYSVTAAKQGIPQVVISRICMA 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 MESKMGELPLDINIQEPRWDQSTPLGRARHFFTVTDPRNLLLSGAQLEASRNIVQNYRAG 60
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Isolated polypoptide with a human transport protein sequence is useful for the diagnosis, prevention and treatment of disorders associated with the immune, reproductive and cardiovascular systems
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VVTPGITEDQLWRAXYVYDSAFHPDTGEKVVLIGRMSAQVPMMMTITGCMLTFYRKTPTV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Duery Match
Best Local Similarity 100.0%; Pred. No. 3.7e-144; Length 374;
Matches 261; Conservative 0; Mismatches 0; Indels 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human ORFX ORF1353 polypeptide sequence SEQ 1D NO:2706.
                                                                                                                                                         Claim 2; Page 108-109; 165pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 374 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43397, which represent the human OSPRX open reading frames 1 to 3161. The OSPRX equences have activities such as: cytostatic; hepatocropic; vulnerary; sequences have activities such as: cytostatic; hemoprotective; vulnerary; osteopathic; antiparkinsonian; nociropic; neuroprotective; vulnerary; osteopathic; antiparkinsonian; nociropic; immunosuppressant; osteopathic; antidabetic; hypotensive; dermatological; immunosuppressant; antidiabetic; hypotensive; dermatological; immunosuppressive; antidiateratic; hypotensive; dermatological; immunosuppressive; antidiateratory; antidiateraris, antitional antithematic; antitional control and antiansemic. The sequences can be used for determining the presence of or predisposition to or preventing or treating pathological conditions ssociated with an ONFX-associated disorder. The moleic acids can be used to express ONFX proteins in gene therapy vectors. The proteins an enclosic acids may be used to treat cancers, or the presence of disorders neuroleic acids may be used to treat cancers, corollerative disorders neuroleic acids may be used to treat cancers, or proliferative disorders neuroleic acids may be used to treat cancers, corollerative disorders neuroleic acids may be used to treat cancers, or proliferation, hypothrodism, cholesacrol ester-storage, systemic lupus of the present of the presence of antidismin the antification, malaria, autoimmune disorders asthma, all all passociated and and artilage damage, cocurrant hasemoglobinuria, antilifiammatory disease; to enhance conductory conductors. The content as a contraceptive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 VVTPGITEDQLWRAKYVYDSAFHPDTGEKVVLIGRMSAQVPWNWTITGCMLTFYRKTPTV 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                            Novel nucleic acids and peptides derived from open reading frame X, useful for treating e.g. cancers, proliferative disorders, neurodegenerative disorders and cardiovascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94.3%; Score 1269; DB 21; Length 251; 98.0%; Pred. No. 9.2e-136; ive 3; Mismatches 2; Indels 0.
                                                                                                                                                                                                                                                                                                                                                       Claim 11; Page 1942-1943; 5507pp; English.
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02-APR-1999; 99US-0127636.
05-APR-1999; 99US-0127728.
30-MAR-2000; 2000US-0540763.
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Best Local Similarity 98.0°
Matches 246; Conservative
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                                                                                                                                           Shimkets RA, Leach M;
                                                                                           (CURA-) CURAGEN CORP.
                                                                                                                                                                                      WPI; 2000-602362/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      251 AA;
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